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<u>L4</u>	L3 same (advantag\$ or useful\$)	6	<u>L4</u>	
<u>L3</u>	L2 same bind\$ same protein	24	<u>L3</u>	
<u>L2</u>	nucleotide near0 analogu\$	904	<u>L2</u>	
<u>L1</u>	nucleotide near0 aalogu\$	0	<u>L1</u>	

END OF SEARCH HISTORY

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=> s l1 (p)bind? (p) (protein or peptide) L2 307 L1 (P) BIND? (P) (PROTEIN OR PEPTIDE)

=> s 12 (p) (advantag? or useful?)
L3 16 L2 (P) (ADVANTAG? OR USEFUL?)

=> duplicate remove 13

DUPLICATE PREFERENCE IS 'MEDLINE, BIOSIS, EMBASE'

KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n

PROCESSING COMPLETED FOR L3

L4 7 DUPLICATE REMOVE L3 (9 DUPLICATES REMOVED)